

GenCore Version 5.1.3
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OM protein - nucleic search, using frame_plus_b2n model

Run on: January 16, 2003, 16:59:22 ; Search time 14:5714 Seconds
(without alignments)
89.447 Million cell updates/sec

Title: US-09-856-070-19

Perfect score: 65

Sequences: 1 KEELEIGHOFF 13

Scoring table:

BLOSUM62
Xgapop 16.0 ; Xgapext 0.5
Ygapop 10.0 ; Ygapext 0.5
Fgapop 6.0 ; Fgapext 7.0
Delop 6.0 ; Deloxt 7.0

Searched: 441362 seqs, 15338381 residues

Total number of hits satisfying chosen parameters: 882724

Minimum DB seq length: 0

Maximum DB seq length: 260000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Command line parameters:

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-Q=/cgn2/6/ptodata/2/ina/5A.COMB.seq.*
-DB=Issued_Patents_NA -OMI=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALLIGN=200 -THR_SCORE=100 -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=tbl -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=nsr0456070@compugen_161@frugal_14012003_15880555 -NPP=6 -LPU 3
-NO_XLPEX -NO_WMAP -LAP=QUERY -NES_SCORES=6 -WAIT -LNAME=DEV_TIMEWIT-120
-WARN_TIMEOUT=30 -THRAUS=1 -XGAPOP=10 -XGAPEXT=0.5 -PGAPOP=6 -FGAPEXT=7
-XGAPOP=10 -YGAPEXT=0.5 -DELLOP=6 -DELEXT=7

Database : Issued_Patents_NA.*

- 1: /cgn2/6/ptodata/2/ina/5A.COMB.seq.*
- 2: /cgn2/6/ptodata/2/ina/5B.COMB.seq.*
- 3: /cgn2/6/ptodata/2/ina/6A.COMB.seq.*
- 4: /cgn2/6/ptodata/2/ina/6B.COMB.seq.*
- 5: /cgn2/6/ptodata/2/ina/PCUS.COMB.seq.*
- 6: /cgn2/6/ptodata/2/ina/backkilled.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	39	60.0	1068	5	PCT-US91-00899-13
2	39	60.0	1215	5	PCT-US91-00899-5
3	39	60.0	1488	1	US-07-914-281-9
4	39	60.0	1488	1	US-08-394-246-4
5	39	60.0	1488	1	US-08-525-058A-9
6	39	60.0	1488	2	US-08-696-731-9
7	39	60.0	1488	4	US-09-042-531-9
8	39	60.0	2134	2	US-08-483-151-3
9	39	60.0	2134	5	PCT-US96-06427-3
10	39	60.0	2175	4	US-08-482-073-9
11	39	60.0	2861	4	US-08-482-073-10
12	39	60.0	1627	4	US-09-323-873A-6

13	39	60.0	3647	1	US-07-914-281-7	Sequence 7, Appl
14	39	60.0	3647	1	US-08-394-246-7	Sequence 7, Appl
15	39	60.0	3647	1	US-08-525-058A-7	Sequence 7, Appl
16	39	60.0	3647	2	US-08-696-731-7	Sequence 7, Appl
17	39	60.0	3647	4	US-09-042-531-7	Sequence 7, Appl
18	39	60.0	3647	5	PCT-US91-00899-4	Sequence 4, Appl
19	38	58.5	1134	4	US-09-206-059-29	Sequence 29, Appl
20	38	58.5	2313	4	US-09-373-838-157	Sequence 157, Appl
21	38	58.5	2497	1	US-08-643-219-12	Sequence 12, Appl
22	38	58.5	2497	2	US-09-131-995-12	Sequence 12, Appl
23	38	58.5	2497	2	US-08-832-087B-12	Sequence 12, Appl
24	38	58.5	2497	3	US-08-851-350-12	Sequence 12, Appl
25	38	58.5	2497	4	US-09-132-154-12	Sequence 12, Appl
26	38	58.5	2753	6	US-09-440-7	Patent No 5200340
27	38	58.5	2753	1	US-07-854-603-1	Sequence 1, Appl
28	38	58.5	28305	4	US-09-453-702B-251	Sequence 251, Appl
29	37	56.9	1969	4	US-09-454-782B-72	Sequence 72, Appl
30	37	56.9	3617	4	US-09-511-057C-12	Sequence 12, Appl
31	37	56.9	3617	4	US-09-513-057C-14	Sequence 14, Appl
32	36.5	56.2	1650	1	US-08-354-240A-1	Sequence 1, Appl
33	36.5	56.2	1650	1	US-08-354-240A-3	Sequence 3, Appl
34	36.5	56.2	1653	1	US-08-354-240A-5	Sequence 5, Appl
35	36.5	56.2	1656	3	US-09-111-752-6	Sequence 6, Appl
36	36.5	56.2	1656	3	US-09-111-752-9	Sequence 9, Appl
37	36.5	56.2	1722	3	US-08-718-425-1	Sequence 1, Appl
38	36.5	56.2	1722	4	US-09-380-061B-1	Sequence 1, Appl
39	36.5	56.2	1725	4	US-09-380-061B-19	Sequence 19, Appl
40	36.5	56.2	1798	4	US-09-446-402A-16	Sequence 16, Appl
41	36.5	56.2	1811	3	US-08-867-352-32	Sequence 32, Appl
42	36.5	56.2	1838	4	US-08-487-183A-9	Sequence 9, Appl
43	36.5	56.2	2445	1	US-08-122-520C-8	Sequence 8, Appl
44	36.5	56.2	4544	4	US-09-488-270A-1	Sequence 1, Appl
45	36.5	56.2	5157	2	US-08-474-169-7	Sequence 7, Appl

ALIGNMENTS

RESULT 1 00899-13
PCT-US91-00899-13

Sequence 13, Application PCT/US91-00899

GENERAL INFORMATION:

APPLICANT: lower, John H.

TITLE OF INVENTION: Method and products for the synthesis of

TITLE OF INVENTION: Oligosaccharide Structures on Glycoproteins, Glycolipids, or as Free Molecules, and for the Isolation of Cloned Genetic Sequences That Determine These Structures

TITLE OF INVENTION: Genetic Sequences That Determine These Structures

NUMBER OF SEQUENCES: 16

CORRESPONDENCE ADDRESS:

ADDRESSER: ORION, SPIVAK, MCCLELLAND, MAIER & NUSSBAUM,

ADDRESSER: P.C.

STREET: 1775 Jefferson Davis Highway, Suite 470

CITY: Arlington

STATE: Virginia

ZIP: 22202

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patent Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US91/00899

FILING DATE: 19910214

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Lavalleye Ph.D., Jean Paul

REGISTRATION NUMBER: 31,451

REFERENCE/BOOK NUMBER: 2363-021-55 PCT

TELEPHONE: (703)521-5940

TELEFAX: (703)486-2347

TELEX: 248855 OPAT UP

INFORMATION FOR SEQ ID NO: 13:

SEQUENCE CHARACTERISTICS:


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: FILING DATE: 30-MAR-1994
: APPLICATION NUMBER: US 07/914,281
: FILING DATE: 20-JUL-1992
: ATTORNEY/AGENT INFORMATION:
: NAME: Lavalleye, Jean-Paul M. P.
: REGISTRATION NUMBER: 31,451
: REFERENCE/DOCKET NUMBER: 2363-060-55
: TELECOMMUNICATION INFORMATION:
: TELEPHONE: (703)521-4500
: TELEFAX: (703)486-2347
: TELEX: 248855 OPAI UR
: INFORMATION FOR SEQ ID NO: 9:
: SEQUENCE CHARACTERISTICS:
: LENGTH: 1488 base pairs
: TYPE: nucleic acid
: STRANDEDNESS: unknown
: TOPOLOGY: unknown
: MOLECULE TYPE: DNA (genomic)
US-08-696-731-9

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Alignment Scores:
Pred. No.: 44.8 Length: 1488
Score: 39.00 Matches: 8
Percent Similarity: 83.33% Conservative: 2
Best Local Similarity: 66.67% Mismatches: 2
Query Match: 60.00% Indels: 0
DB: 2 Gaps: 0

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US 09-856-070-19 (1-13) x US 08-696-731-9 (1-1488)

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QY 2 GluGluMetLeuArgLeuGlnAspTyrGluGlu 13
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DB 679 GAGGAGGTGGATCGAGGAGGAGGAGGAGGAG 714

```

RESULT 7

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US 09-042-531-9
: Sequence 9, Application US/09042531
: Patent No. 6268193
: GENERAL INFORMATION:
: APPLICANT: LOWE, JOHN R.
: TITLE OF INVENTION: METHODS AND PRODUCTS FOR THE SYNTHESIS
: TITLE OF INVENTION: OF OLIGOSACCHARIDE STRUCTURES ON GLYCOPROTEINS,
: TITLE OF INVENTION: GLYCOLIPIDS, OR AS FREE MOLECULES, AND FOR THE ISOLATION
: TITLE OF INVENTION: OF CLONED GENETIC SEQUENCES THAT DETERMINE THESE STRUCTU
: NUMBER OF SEQUENCES: 14
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: OHION, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
: STREET: 1755 Jefferson Davis Highway, Fourth Floor
: CITY: Arlington
: STATE: Virginia
: COUNTRY: U.S.A.
: ZIP: 22202
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: Patent In Release #1.0, Version #1.25
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/09/042 531
: FILING DATE:
: CLASSIFICATION:
: PRIOR APPLICATION DATA:
: APPLICATION NUMBER: US/09/042 531
: FILING DATE:
: APPLICATION NUMBER: US 07/220,433
: FILING DATE: 30-MAR-1994
: APPLICATION NUMBER: US 07/914,281
: FILING DATE: 20-JUL-1992
: ATTORNEY/AGENT INFORMATION:
: NAME: Lavalleye, Jean-Paul M. P.
: REGISTRATION NUMBER: 31,451
: REFERENCE/DOCKET NUMBER: 2363-060-55

```

```

: TELECOMMUNICATION INFORMATION:
: TELEPHONE: (703)521-4500
: TELEFAX: (703)486-2347
: TELEX: 248855 OPAI UR
: INFORMATION FOR SEQ ID NO: 9:
: SEQUENCE CHARACTERISTICS:
: LENGTH: 1488 base pairs
: TYPE: nucleic acid
: STRANDEDNESS: unknown
: TOPOLOGY: unknown
: MOLECULE TYPE: DNA (genomic)
US-09-042-531-9

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Alignment Scores:
Pred. No.: 44.8 Length: 1488
Score: 39.00 Matches: 8
Percent Similarity: 83.33% Conservative: 2
Best Local Similarity: 66.67% Mismatches: 2
Query Match: 60.00% Indels: 0
DB: 2 Gaps: 0

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US 09-856-070-19 (1-13) x US-09-042-531-9 (1-1488)

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QY 2 GluGluMetLeuArgLeuGlnAspTyrGluGlu 14
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DB 675 GAGGAGGTGGATCGAGGAGGAGGAGGAG 714

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RESULT 8

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US-08-483-151-3
: Sequence 3, Application US/08483151
: Patent No. 5858752
: GENERAL INFORMATION:
: APPLICANT: Seod, Brian
: TITLE OF INVENTION: FLYCUT TRANSFERPASP GFNPS AND USVS THEREOF
: NUMBER OF SEQUENCES: 4
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: Fish & Richardson P.C.
: STREET: 225 Franklin Street
: CITY: Boston
: STATE: MA
: COUNTRY: USA
: ZIP: 02110-2804
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: Patent In Release #1.0, Version #1.30
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/08/483,151
: FILING DATE: 07-JUN-1995
: CLASSIFICATION: 530
: ATTORNEY/AGENT INFORMATION:
: NAME: Lech, Karen F.
: REGISTRATION NUMBER: 35,238
: REFERENCE/DOCKET NUMBER: 00/86/278001
: TELECOMMUNICATION INFORMATION:
: TELEPHONE: 617/542-5070
: TELEFAX: 617/542-8906
: TELEX: 200154
: INFORMATION FOR SEQ ID NO: 3:
: SEQUENCE CHARACTERISTICS:
: LENGTH: 2134 base pairs
: TYPE: nucleic acid
: STRANDEDNESS: single
: TOPOLOGY: linear
: MOLECULE TYPE: cDNA
US-08-483-151-3

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```

Alignment Scores:
Pred. No.: 68.9 Length: 2134
Score: 39.00 Matches: 8
Percent Similarity: 83.33% Conservative: 2

```



```

RESULT 11
US 08 482-073 10
? Sequence 10, Application US/08482073
? Patent No. 6407025
? GENERAL INFORMATION:
? APPLICANT: Hession, Catherine A.
? APPLICANT: Lobb, Roy R.
? APPLICANT: Goetz, Susan E.
? APPLICANT: Osborn, Laurelee
? APPLICANT: Benjamin, Christopher D.
? APPLICANT: Kosa, Margaret D.
? TITLE OF INVENTION: ENDOTHELIAL CELL-LEUKOCYTE ADHESION
? TITLE OF INVENTION: MOLECULES (ELAMS) AND MOLECULES INVOLVED IN LEUKOCYTE
? TITLE OF INVENTION: ADHESION (WILAS)
? NUMBER OF SEQUENCES: 25
? CORRESPONDENCE ADDRESS:
? ADDRESSEE: Fish & Neave
? STREET: 1251 Avenue of the Americas
? CITY: New York
? STATE: New York
? COUNTRY: United States of America
? ZIP: 10020
? COMPUTER READABLE FORM:
? MEDIUM TYPE: Floppy disk
? COMPUTER: IBM PC compatible
? OPERATING SYSTEM: PC-DOS/MS-DOS
? SOFTWARE: Patent In Release #1.0, Version #1.25
? CURRENT APPLICATION DATA:
? APPLICATION NUMBER: US/08/482,073
? FILING DATE:
? CLASSIFICATION:
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US/08/486,336
? FILING DATE:
? APPLICATION NUMBER: US 07/608298
? FILING DATE: 31-OCT-1990
? PRIOR APPLICATION NUMBER: WO PCT/US 90/02357
? FILING DATE: 27-APR-1990
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US 07/452675
? FILING DATE: 18-DEC-1989
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US 07/354516
? FILING DATE: 01-JUN-1989
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US 07/354151
? FILING DATE: 28-APR-1989
? ATTORNEY/AGENT INFORMATION:
? NAME: Haley Jr., James F.
? REGISTRATION NUMBER: 27,794
? REFERENCE/DOCKET NUMBER: B124CIP4
? TELECOMMUNICATION INFORMATION:
? TELEPHONE: (212) 596-9000
? TELEFAX: (212) 596-9090
? TELEX: 14-8167
? INFORMATION FOR SEQ ID NO: 10:
? SEQUENCE CHARACTERISTICS:
? LENGTH: 2861 base pairs
? TYPE: nucleic acid
? STRANDEDNESS: double
? TOPOLOGY: linear
US-08-482-073-10

Alignment Scores:
? Aligned No. 47 4 Length: 2861
? Score: 39.00 Matches: 8
? Percent Similarity: 83.33% Conservative: 2
? Best Local Similarity: 66.67% Mismatches: 2
? Query Match: 60.00% Indels: 0
? DB: 4 Gaps: 0
US-08-482-070-19 (1-13) x US-08-482-073-10 (1-2861)

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FILING DATE: 19920720
 CLASSIFICATION: 530
 ATTORNEY/AGENT INFORMATION:
 NAME: Lavalleye, Jean-Paul M. P.
 REGISTRATION NUMBER: 31,451
 REFERENCE/DOCKET NUMBER: 2363-060-55
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (703)521-4500
 TELEFAX: (703)486-2347
 INFORMATION FOR SEQ ID NO: 7:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 3647 base pairs
 TYPE: NUCLEIC ACID
 STRANDEDNESS: unknown
 TOPOLOGY: unknown
 MOLECULE TYPE: DNA (genomic)
 ANTI-SENSE: NO
 US-07-914-281-7

Alignment Scores:
 Pred. No.: 131 Length: 3647
 Score: 39.00 Matches: 8
 Percent Similarity: 83.33% Conservative: 2
 Best Local Similarity: 66.67% Mismatches: 2
 Query Match: 60.00% Indels: 0
 DB: 1 Gaps: 0

US-09-856-070-19 (1-13) x US 07 914 281 7 (1-3647)

QY 2 GluGluLeuMetLeuArgLeuGluAspTyrGluGln 13
 DB 2347 GAGGAGGCGGAGTGTGGCGCTGTCACATACAGAG 2392

RESULT 14

US-08-393-246-7

Sequence 7, Application US/08193246

Patent No. 5595900

GENERAL INFORMATION:

APPLICANT: LOWE, JOHN B.

TITLE OF INVENTION: METHODS AND PRODUCTS FOR THE SYNTHESIS

TITLE OF INVENTION: OF OLIGOSACCHARIDE STRUCTURES ON GLYCOPROTEINS,

TITLE OF INVENTION: GLYCOLIPIDS, OR AS FREE MOLECULES, AND FOR THE ISOLATION

TITLE OF INVENTION: OF CLONED GENETIC SEQUENCES THAT DETERMINE THESE STRUCTU

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:

ADDRESSEE: OHION, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,

ADDRESS: P.C.

STREET: 1755 Jefferson Davis Highway, Fourth Floor

CITY: Arlington

STATE: Virginia

COUNTRY: U.S.A.

ZIP: 22202

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/393,246

FILING DATE:

CLASSIFICATION: 530

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/220,433

FILING DATE: 30-MAR-1994

APPLICATION NUMBER: US 07/914,281

FILING DATE: 20-JUL-1992

ATTORNEY/AGENT INFORMATION:

NAME: Lavalleye, Jean-Paul M. P.

REGISTRATION NUMBER: 31,451

REFERENCE/DOCKET NUMBER: 2363-060-55

TELECOMMUNICATION INFORMATION:

TELEPHONE: (703)521-4500

TELEFAX: (703)486-2347
 TELEPHONE: 248855 OPAT UR
 INFORMATION FOR SEQ ID NO: 7:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 3647 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: unknown
 TOPOLOGY: unknown
 MOLECULE TYPE: DNA (genomic)
 ANTI-SENSE: NO
 US-08-393-246-7

Alignment Scores:
 Pred. No.: 131 Length: 3647
 Score: 39.00 Matches: 8
 Percent Similarity: 83.33% Conservative: 2
 Best Local Similarity: 66.67% Mismatches: 2
 Query Match: 60.00% Indels: 0
 DB: 1 Gaps: 0

US-09-856-070-19 (1-13) x US-08-393-246-7 (1-3647)

QY 2 GluGluLeuMetLeuArgLeuGluAspTyrGluGln 13

DB 2347 GAGGAGGCGGAGTGTGGCGCTGTCACATACAGAG 2392

RESULT 15

US-08-525-058A-7

Sequence 7, Application US/08525058A

Patent No. 5770420

GENERAL INFORMATION:

APPLICANT: LOWE, JOHN B.

TITLE OF INVENTION: METHODS AND PRODUCTS FOR THE SYNTHESIS

TITLE OF INVENTION: OF OLIGOSACCHARIDE STRUCTURES ON GLYCOPROTEINS,

TITLE OF INVENTION: GLYCOLIPIDS, OR AS FREE MOLECULES, AND FOR THE ISOLATION

TITLE OF INVENTION: OF CLONED GENETIC SEQUENCES THAT DETERMINE THESE STRUCTURES

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:

ADDRESSEE: OHION, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C.

ADDRESS: 1755 Jefferson Davis Highway, Fourth Floor

CITY: Arlington

STATE: Virginia

COUNTRY: U.S.A.

ZIP: 22202

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/525,058A

FILING DATE:

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Lavalleye, Jean-Paul M. P.

REGISTRATION NUMBER: 31,451

REFERENCE/DOCKET NUMBER: 2363-060-55

TELECOMMUNICATION INFORMATION:

TELEPHONE: (703)521-4500

TELEFAX: (703)486-2347

INFORMATION FOR SEQ ID NO: 7:

SEQUENCE CHARACTERISTICS:

LENGTH: 3647 base pairs

TYPE: nucleic acid

STRANDEDNESS: double

TOPOLOGY: linear

MOLECULE TYPE: DNA (genomic)

ANTI-SENSE: NO

US-08-525-058A-7

Alignment Scores:

Pred. No.: 131 Length: 3647

